



PCT10

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/070,588

DATE: 01/30/2003

TIME: 10:33:27

Input Set : A:\EP.txt

Output Set: N:\CRF4\01302003\J070588.raw

4 <110> APPLICANT: EPIDAUROS AG
 6 <120> TITLE OF INVENTION: Polymorphisms in the human hPXR gene and their use in
 7 diagnostic and therapeutic applications
 9 <130> FILE REFERENCE: D 2145 PCT-2
 OK-> 11 <140> CURRENT APPLICATION NUMBER: US/10/070,588
 OK-> 12 <141> CURRENT FILING DATE: 2002-10-15
 14 <160> NUMBER OF SEQ ID NOS: 185
 16 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

2706 <210> SEQ ID NO: 185
 2707 <211> LENGTH: 324
 2708 <212> TYPE: DNA
 2709 <213> ORGANISM: Homo sapiens
 2711 <220> FEATURE:
 2712 <223> OTHER INFORMATION: r=g or a
 2714 <400> SEQUENCE: 185
 2715 tgcttggtgca gcctcagagc agccctgagg cttgtgggtc agggcgggct gcaccacaaa 60
 2716 tcttttctct ggctggcatg caggttcttg ttctgaaga tcatggctat gtcaccgag 120
 2717 ctccgcagca tcaatgctca gcacaccag cggtgctgc gcatccagga catacacccc 180
 2718 tttgctacgc ccctcatgca ggagttgttc ggcatcacag gtagctgagc rgctgccctt 240
 2719 ggrrgacacc tccgagaggc agccagacc agagccctct gagccgccac tcccgggcca 300
 2720 agacagatgg aactgcca gagc 324

Does Not Comply
Corrected Diskette Needed

pp 1-3

E--> 2723 1

E--> 2726 43

E--> 2741 76

delete

see p. 2 for more errors

<210> 1
<211> 18
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Artificial
sequence

<400> 1

tcatgtccga cgaggccg

(global error)

insufficient explanation -
give source
of genetic
material

18

The types of errors shown exist throughout
the Sequence Listing. Please check subsequent
sequences for similar errors.

(see item 11
on Error
Summary Sheet)

11 Sequence: error

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/10/070,588

DATE: 01/30/2003

TIME: 10:33:28

Input Set : A:\EP.txt

Output Set: N:\CRF4\01302003\J070588.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.

Use of <220> to <223> is MANDATORY if n's or Xaa's are present.

in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:178; N Pos. 780.

VERIFICATION SUMMARY

PATENT APPLICATION: **US/10/070,588**

DATE: 01/30/2003

TIME: 10:33:28

Input Set : **A:\EP.txt**

Output Set: **N:\CRF4\01302003\J070588.raw**

L:11 M:270 C: Current Application Number differs, Replaced Application Number
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:2586 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:178
L:2586 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:178
L:2586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:178 after pos.:720
L:2723 M:254 E: No. of Bases conflict, this line has no nucleotides.
M:254 Repeated in SeqNo=185